



#14

## SEQUENCE LISTING

<110> Ono Pharmaceutical Co., Ltd.

<120> NOVEL PLASMID DNA COMPRISING REPORTER GENE DNA AND USE OF THE SAME

<130> Q57282

<140> 09/446,634

<141> 1999-12-23

<150> JP 9-171440

<151> 1997-06-27

<160> 23

<170> PatentIn version 3.2

<210> 1

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer F1

<400> 1  
ccaagcttgg cgaccagcaa tacaaactgc aggaaac 37

<210> 2

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer R1

<400> 2  
tcaggatcca gacattgtcc ttcattttca tt 32

<210> 3

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer F2

<400> 3  
aaccagcacc atctgggtgc gatggt 26

<210> 4  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer R2

<400> 4  
aggtgtggct gatctgaagg aactca

26

<210> 5  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer F3

<400> 5  
agaaatgacc atggttgaca cagaga

26

<210> 6  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer R3

<400> 6  
aaatgttggc agtggctcag gactct

26

<210> 7  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer F4

<400> 7  
agatcagcca tggagcagcc acagga

26

<210> 8  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer R4

<400> 8  
 attggagtct gcagggaggc ctgggt 26

<210> 9  
 <211> 37  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer F5

<400> 9  
 gcaagcttca ccatgaagct actgtcttct atcgaac 37

<210> 10  
 <211> 33  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer R5

<400> 10  
 agccatggcc ggcgatacag tcaactgtct ttg 33

<210> 11  
 <211> 62  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer F6

<400> 11  
 gccatggctc ctaagaagaa gcgtaaggta ggatcccata atgccatcag gtttgggagg 60

at 62

<210> 12  
 <211> 69  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer R6

<400> 12  
 cctctagact agctggcata gtcgggcacg tcgtaggggt agtcgacgta caagtccttg 60

tagatctcc 69

<210> 13  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer F7

<400> 13  
cacggatccc acaacgcgat tcgttttgga cga 33

<210> 14  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer R7

<400> 14  
atggtcgacg tacatgtccc ttagatctc ctg 33

<210> 15  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer F8

<400> 15  
cacggatccc acaacgctat ccgttttggt cgg 33

<210> 16  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer R8

<400> 16  
atggtcgacg tacatgtcct ttagatctc ctg 33

<210> 17  
<211> 38  
<212> DNA  
<213> Artificial Sequence

<220>  
 <223> Multiple cloning site DNA linker - sense strand  
  
 <400> 17  
 gaattcgtcg acggtaccga tatcgagctc gcggccgc 38  
  
 <210> 18  
 <211> 85  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Gal4 responsive element  
  
 <400> 18  
 tcgacggagt actgtcctcc gcgacggagt actgtcctcc gcgacggagt actgtcctcc 60  
 gcgacggagt actgtcctcc gagct 85  
  
 <210> 19  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Gal4 responsive element  
  
 <400> 19  
 cgacggagta ctgtcctccg 20  
  
 <210> 20  
 <211> 9  
 <212> PRT  
 <213> Artificial Sequence  
  
 <220>  
 <223> Influenza hemagglutinin epitope  
  
 <400> 20  
  
 Tyr Pro Tyr Asp Val Pro Asp Tyr Ala  
 1 5  
  
 <210> 21  
 <211> 38  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Multiple cloning site DNA linker - antisense strand

<400> 21  
gcggccgcga gctcgatata ggtaccgtcg acgaattc

38

<210> 22  
<211> 335  
<212> PRT  
<213> Homo sapiens

<220>  
<221> mat\_peptide  
<222> (17)..(355)

<400> 22

Met Leu Gly Ile Trp Thr Leu Leu Pro Leu Val Leu Thr Ser Val Ala  
-15 -10 -5 -1

Arg Leu Ser Ser Lys Ser Val Asn Ala Gln Val Thr Asp Ile Asn Ser  
1 5 10 15

Lys Gly Leu Glu Leu Arg Lys Thr Val Thr Thr Val Glu Thr Gln Asn  
20 25 30

Leu Glu Gly Leu His His Asp Gly Gln Phe Cys His Lys Pro Cys Pro  
35 40 45

Pro Gly Glu Arg Lys Ala Arg Asp Cys Thr Val Asn Gly Asp Glu Pro  
50 55 60

Asp Cys Val Pro Cys Gln Glu Gly Lys Glu Tyr Thr Asp Lys Ala His  
65 70 75 80

Phe Ser Ser Lys Cys Arg Arg Cys Arg Leu Cys Asp Glu Gly His Gly  
85 90 95

Leu Glu Val Glu Ile Asn Cys Thr Arg Thr Gln Asn Thr Lys Cys Arg  
100 105 110

Cys Lys Pro Asn Phe Phe Cys Asn Ser Thr Val Cys Glu His Cys Asp  
115 120 125

Pro Cys Thr Lys Cys Glu His Gly Ile Ile Lys Glu Cys Thr Leu Thr  
130 135 140

Ser Asn Thr Lys Cys Lys Glu Glu Gly Ser Arg Ser Asn Leu Gly Trp  
 145 150 155 160

Leu Cys Leu Leu Leu Leu Pro Ile Pro Leu Ile Val Trp Val Lys Arg  
 165 170 175

Lys Glu Val Gln Lys Thr Cys Arg Lys His Arg Lys Glu Asn Gln Gly  
 180 185 190

Ser His Glu Ser Pro Thr Leu Asn Pro Glu Thr Val Ala Ile Asn Leu  
 195 200 205

Ser Asp Val Asp Leu Ser Lys Tyr Ile Thr Thr Ile Ala Gly Val Met  
 210 215 220

Thr Leu Ser Gln Val Lys Gly Phe Val Arg Lys Asn Gly Val Asn Glu  
 225 230 235 240

Ala Lys Ile Asp Glu Ile Lys Asn Asp Asn Val Gln Asp Thr Ala Glu  
 245 250 255

Gln Lys Val Gln Leu Leu Arg Asn Trp His Gln Leu His Gly Lys Lys  
 260 265 270

Glu Ala Tyr Asp Thr Leu Ile Lys Asp Leu Lys Lys Ala Asn Leu Cys  
 275 280 285

Thr Leu Ala Glu Lys Ile Gln Thr Ile Ile Leu Lys Asp Ile Thr Ser  
 290 295 300

Asp Ser Glu Asn Ser Asn Phe Arg Asn Glu Ile Gln Ser Leu Val  
 305 310 315

<210> 23  
 <211> 327  
 <212> PRT  
 <213> Mus musculus

<220>  
 <221> mat\_peptide  
 <222> (22)..(327)

<400> 23

Met Leu Trp Ile Trp Ala Val Leu Pro Leu Val Leu Ala Gly Ser Gln  
-20 -15 -10

Leu Arg Val His Thr Gln Gly Thr Asn Ser Ile Ser Glu Ser Leu Lys  
-5 -1 1 5 10

Leu Arg Arg Arg Val His Glu Thr Asp Lys Asn Cys Ser Glu Gly Leu  
15 20 25

Tyr Gln Gly Gly Pro Phe Cys Cys Gln Pro Cys Gln Pro Gly Lys Lys  
30 35 40

Lys Val Glu Asp Cys Lys Met Asn Gly Gly Thr Pro Thr Cys Ala Pro  
45 50 55

Cys Thr Glu Gly Lys Glu Tyr Met Asp Lys Asn His Tyr Ala Asp Lys  
60 65 70 75

Cys Arg Arg Cys Thr Leu Cys Asp Glu Glu His Gly Leu Glu Val Glu  
80 85 90

Thr Asn Cys Thr Leu Thr Gln Asn Thr Lys Cys Lys Cys Lys Pro Asp  
95 100 105

Phe Tyr Cys Asp Ser Pro Gly Cys Glu His Cys Val Arg Cys Ala Ser  
110 115 120

Cys Glu His Gly Thr Leu Glu Pro Cys Thr Ala Thr Ser Asn Thr Asn  
125 130 135

Cys Arg Lys Gln Ser Pro Arg Asn Arg Leu Trp Leu Leu Thr Ile Leu  
140 145 150 155

Val Leu Leu Ile Pro Leu Val Phe Ile Tyr Arg Lys Tyr Arg Lys Arg  
160 165 170

Lys Cys Trp Lys Arg Arg Gln Asp Asp Pro Glu Ser Arg Thr Ser Ser  
175 180 185

Arg Glu Thr Ile Pro Met Asn Ala Ser Asn Leu Ser Leu Ser Lys Tyr



190	195	200
Ile Pro Arg Ile Ala Glu Asp Met Thr Ile Gln Glu Ala Lys Lys Phe		
205	210	215
Ala Arg Glu Asn Asn Ile Lys Glu Gly Lys Ile Asp Glu Ile Met His		
220	225	230
Asp Ser Ile Gln Asp Thr Ala Glu Gln Lys Val Gln Leu Leu Leu Cys		
	240	245
Trp Tyr Gln Ser His Gly Lys Ser Asp Ala Tyr Gln Asp Leu Ile Lys		
	255	260
Gly Leu Lys Lys Ala Glu Cys Arg Arg Thr Leu Asp Lys Phe Gln Asp		
	270	275
Met Val Gln Lys Asp Leu Gly Lys Ser Thr Pro Asp Thr Gly Asn Glu		
	285	290
Asn Glu Gly Gln Cys Leu Glu		
300	305	